

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 09/804,733B

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000
- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.



1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/804,733B

DATE: 12/18/2002

TIME: 12:01:09

Input Set : A:\MONS016US.TXT

Output Set: N:\CRF4\12182002\I804733B.raw

2 <110> APPLICANT: WANG et al.
 3 <120> TITLE OF INVENTION: RECOMBINANT PROTEINS CONTAINING REPEATING UNITS
 4 <130> FILE REFERENCE: MONS:016US
 5 <140> CURRENT APPLICATION NUMBER: 09/804,733B
 6 <141> CURRENT FILING DATE: 2001-03-13
 7 <150> PRIOR APPLICATION NUMBER: 60/188,990
 8 <151> PRIOR FILING DATE: 2000-03-13
 9 <160> NUMBER OF SEQ ID NOS: 29
 10 <170> SOFTWARE: PatentIn version 3.0
 11 <110> SEQ ID NO: 1
 12 <111> LENGTH: 5
 13 <112> TYPE: PRT
 14 <113> ORGANISM: Euthyrnus pelamis
 15 <114> FEATURE:
 16 <115> NAME/KEY: PEPTIDE
 17 <116> LOCATION: (1)..(5)
 18 <400> SEQUENCE: 1
 19 Lys Pro Asn Met
 20 1 5
 21 <210> SEQ ID NO: 2
 22 <211> LENGTH: 4
 23 <212> TYPE: PRT
 24 <213> ORGANISM: Euthyrnus pelamis
 25 <214> FEATURE:
 26 <215> NAME/KEY: PEPTIDE
 27 <216> LOCATION: (1)..(4)
 28 <400> SEQUENCE: 2
 29 Lys Pro Asn Met
 30 1
 31 <310> SEQ ID NO: 3
 32 <311> LENGTH: 4
 33 <312> TYPE: PRT
 34 <313> ORGANISM: Euthyrnus pelamis
 35 <314> FEATURE:
 36 <315> NAME/KEY: PEPTIDE
 37 <316> LOCATION: (1)..(4)
 38 <400> SEQUENCE: 3
 39 Val Val Tyr Pro
 40 1
 41 <410> SEQ ID NO: 4
 42 <411> LENGTH: 15
 43 <412> TYPE: DNA
 44 <213> ORGANISM: Artificial/Unknown

C--> 64 <213> ORGANISM: Artificial/Unknown

Does Not Comply
Corrected Diskette Needed

pg 1-6

invalid response - see item 10 on
Error Summary
sheet

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/804,733B

DATE: 11-14-2009

TIME: 10:01:14

Input Set : A:\MONS016US.TXT

Output Set: N:\CRF4\12182002\I804733B.raw

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66 <220> FEATURE:
67 <221> NAME/KEY: misc_feature
68 <222> LOCATION: (1)..(15)
69 <23> OTHER INFORMATION: Degenerate sequence
72 <220> FEATURE:
73 <221> NAME/KEY: misc_feature
74 <222> LOCATION: (1)..(15)
75 <23> OTHER INFORMATION: n=a, t, c or g; r=a or g; y=c or t
76 <400> SEQUENCE: 4
79 ctnaarccna ayatg 15
82 <210> SEQ ID NO: 5
83 <211> LENGTH: 60
84 <212> TYPE: DNA
C--> 85 <213> ORGANISM: Artificial/Unknown
87 <220> FEATURE:
88 <221> NAME/KEY: misc_feature
89 <222> LOCATION: (1)..(60)
90 <23> OTHER INFORMATION: n=any nucleotide; r=a or g; y=c or t
93 <220> FEATURE:
94 <221> NAME/KEY: misc_feature
95 <222> LOCATION: (1)..(60)
96 <23> OTHER INFORMATION: Degenerate sequence
99 <400> SEQUENCE: 5
100 ctnaarccna ayatgctnaa rcnaayatg ctnaarccna ayatgctnaa rcnaayatg 60
103 <210> SEQ ID NO: 6
104 <211> LENGTH: 60
105 <212> TYPE: DNA
C--> 106 <213> ORGANISM: Artificial/Unknown
108 <220> FEATURE:
109 <221> NAME/KEY: misc_feature
110 <222> LOCATION: (1)..(60)
111 <23> OTHER INFORMATION: n=any nucleotide, r=a or g, y=c or t
114 <220> FEATURE:
115 <221> NAME/KEY: misc_feature
116 <222> LOCATION: (1)..(60)
117 <23> OTHER INFORMATION: degenerate sequence
120 <400> SEQUENCE: 6
121 catrttnggy ttnagcatrt tnggyttnag catrttnggy ttnagcatrt tnggyttnag 60
124 <210> SEQ ID NO: 7
125 <211> LENGTH: 25
126 <212> TYPE: DNA
C--> 127 <213> ORGANISM: Artificial/Unknown
129 <220> FEATURE:
130 <221> NAME/KEY: misc_feature
131 <222> LOCATION: (1)..(25)
132 <23> OTHER INFORMATION: Primer
135 <220> FEATURE:
136 <221> NAME/KEY: misc_feature
137 <222> LOCATION: (1)..(25)

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/804,733B

DATE: 12-17-2002

TIME: 12:1:52

Input Set : A:\MONS016US.TXT

Output Set: N:\CRF4\12182002\I804733B.raw

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135 <215> OTHER INFORMATION: n=any nucleotide; r=a or g; y=c or t
141 <410> SEQUENCE: 7
142 aaagaattcc tnaarccnaa yatgc
145 <210> SEQ ID NO: 8
146 <211> LENGTH: 17
147 <212> TYPE: DNA
C--> 148 <213> ORGANISM: Artificial/Unknown
150 <220> FEATURE:
151 <221> NAME/KEY: misc_feature
152 <222> LOCATION: (1)..(17)
153 <223> OTHER INFORMATION: Primer
154 <220> FEATURE:
155 <221> NAME/KEY: misc_feature
156 <222> LOCATION: (1)..(17)
157 <223> OTHER INFORMATION: n=any nucleotide; r=a or g; y=c or t
161 <410> SEQUENCE: 8
163 aaagcgcccg ccatrttngg yttnagc
166 <210> SEQ ID NO: 9
167 <211> LENGTH: 10
168 <212> TYPE: DNA
C--> 169 <213> ORGANISM: Artificial/Unknown
171 <220> FEATURE:
172 <221> NAME/KEY: misc_feature
173 <222> LOCATION: (1)..(10)
174 <223> OTHER INFORMATION: Primer
175 <410> SEQUENCE: 9
176 aaatccggt cactataggg
181 <210> SEQ ID NO: 10
182 <211> LENGTH: 19
183 <212> TYPE: DNA
C--> 184 <213> ORGANISM: Artificial/Unknown
186 <220> FEATURE:
187 <221> NAME/KEY: misc_feature
188 <222> LOCATION: (1)..(19)
189 <223> OTHER INFORMATION: Primer
190 <410> SEQUENCE: 10
191 aaataaatia cgaatgqcc
196 <210> SEQ ID NO: 11
197 <211> LENGTH: 48
198 <212> TYPE: DNA
C--> 199 <213> ORGANISM: Artificial/Unknown
201 <220> FEATURE:
202 <221> NAME/KEY: misc_feature
203 <222> LOCATION: (1)..(48)
204 <223> OTHER INFORMATION: n=any nucleotide; y=c or t
207 <220> FEATURE:
208 <221> NAME/KEY: misc_feature
209 <222> LOCATION: (1)..(48)
210 <223> OTHER INFORMATION: Degenerate sequence

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/804,733B

DATE: 11/15/02

TIME: 10:11:14

Input Set : A:\MONS016US.TXT

Output Set: N:\CRF4\12182002\I804733B.raw

213 <400> SEQUENCE: 11
 214 gtngtntayc cngtngtnta yccngtngtn taycngtng tntayccn 48
 217 <110> SEQ ID NO: 12
 218 <111> LENGTH: 48
 219 <112> TYPE: DNA
 C--> 220 <213> ORGANISM: Artificial/Unknown
 222 <220> FEATURE:
 223 <221> NAME/KEY: misc_feature
 224 <222> LOCATION: (1)..(48)
 225 <223> OTHER INFORMATION: n=any nucleotide; r=a or g
 226 <224> FEATURE:
 227 <221> NAME/KEY: misc_feature
 228 <222> LOCATION: (1)..(48)
 229 <223> OTHER INFORMATION: Degenerate sequence
 234 <400> SEQUENCE: 12
 235 nggrtanacn acnggrtana cnaenggrta nacnaenggr tanacnac 48
 238 <210> SEQ ID NO: 13
 239 <211> LENGTH: 33
 240 <212> TYPE: DNA
 C--> 241 <213> ORGANISM: Artificial/Unknown
 243 <220> FEATURE:
 244 <221> NAME/KEY: misc_feature
 245 <222> LOCATION: (1)..(33)
 246 <223> OTHER INFORMATION: Forward primer
 247 <224> FEATURE:
 248 <221> NAME/KEY: misc_feature
 249 <222> LOCATION: (1)..(33)
 250 <223> OTHER INFORMATION: n=any nucleotide; y=c or t
 254 <400> SEQUENCE: 13
 256 aaaggatccg tngtntaycc ngtngtntay ccn 33
 259 <210> SEQ ID NO: 14
 260 <211> LENGTH: 30
 261 <212> TYPE: DNA
 C--> 262 <213> ORGANISM: Artificial/Unknown
 264 <220> FEATURE:
 265 <221> NAME/KEY: misc_feature
 266 <222> LOCATION: (1)..(30)
 267 <223> OTHER INFORMATION: Reverse primer
 268 <224> FEATURE:
 269 <221> NAME/KEY: misc_feature
 270 <222> LOCATION: (1)..(30)
 271 <223> OTHER INFORMATION: n=any nucleotide; r=a or g
 276 <400> SEQUENCE: 14
 277 cccaagcttn ggrrtanacna cnggrtanac nac 33
 280 <210> SEQ ID NO: 15
 281 <211> LENGTH: 40
 282 <212> TYPE: DNA
 C--> 283 <213> ORGANISM: Artificial/Unknown
 285 <220> FEATURE:

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/804,733B

DATE: 12-17-2002

TIME: 14:11:14

Input Seq : A:\MONS016US.TXT

Output Seq : N:\CRF4\12182002\I804733B.raw

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286 <213> NAME/KEY: misc_feature
287 <213> LOCATION: (1)..(45)
288 <213> OTHER INFORMATION: n=any nucleotide
289 <213> FEATURE:
290 <213> NAME/KEY: misc_feature
291 <213> LOCATION: (1)..(45)
292 <213> OTHER INFORMATION: Degenerate sequence
293 <213> SEQUENCE: 15
OK--> 298 gtnccncng tncncncngt nccncngtn ccncngtnc cncnc 45
301 <213> SEQ ID NO: 16
302 <213> LENGTH: 45
303 <213> TYPE: DNA
C--> 304 <213> ORGANISM: Artificial/Unknown
305 <213> FEATURE:
306 <213> NAME/KEY: misc_feature
307 <213> LOCATION: (1)..(45)
308 <213> OTHER INFORMATION: n=any nucleotide
309 <213> FEATURE:
310 <213> NAME/KEY: misc_feature
311 <213> LOCATION: (1)..(45)
312 <213> OTHER INFORMATION: Degenerate sequence
313 <213> SEQUENCE: 16
OK--> 319 ngngngnacn ggnggnacng gnggnacngg nggnacnggn ggnac 45
322 <213> SEQ ID NO: 17
323 <213> LENGTH: 36
324 <213> TYPE: DNA
C--> 325 <213> ORGANISM: Artificial/Unknown
326 <213> FEATURE:
327 <213> NAME/KEY: misc_feature
328 <213> LOCATION: (1)..(36)
329 <213> OTHER INFORMATION: Forward primer
330 <213> FEATURE:
331 <213> NAME/KEY: misc_feature
332 <213> LOCATION: (1)..(36)
333 <213> OTHER INFORMATION: n=any nucleotide
334 <213> SEQUENCE: 17
OK--> 340 aaagatccg tncncncgt nccncngtn ccncnc 36
341 <213> SEQ ID NO: 18
342 <213> LENGTH: 36
343 <213> TYPE: DNA
C--> 346 <213> ORGANISM: Artificial/Unknown
347 <213> FEATURE:
348 <213> NAME/KEY: misc_feature
349 <213> LOCATION: (1)..(36)
350 <213> OTHER INFORMATION: Reverse primer
351 <213> FEATURE:
352 <213> NAME/KEY: misc_feature
353 <213> LOCATION: (1)..(36)
354 <213> OTHER INFORMATION: n=any nucleotide

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... split throughout
 ... work was done
 ...

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/804,733B

DATE: 12/18/02
TIME: 11:11

Input Set : A:\MONS016US.TXT
Output Set: N:\CRF4\12182002\I804733B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:4; N Pos. 3,9
Seq#:5; N Pos. 3,9,18,24,33,39,48,54
Seq#:6; N Pos. 7,13,22,28,37,43,52,58
Seq#:7; N Pos. 12,18
Seq#:8; N Pos. 18,24
Seq#:11; N Pos. 3,6,12,15,18,24,27,30,36,39,42,48
Seq#:12; N Pos. 1,7,10,13,19,22,25,31,34,37,43,46
Seq#:13; N Pos. 12,15,21,24,27,33
Seq#:14; N Pos. 10,16,19,22,28,31
Seq#:15; N Pos. 3,6,9,12,15,18,21,24,27,30,33,36,39,42,45
Seq#:16; N Pos. 1,4,7,10,13,16,19,22,25,28,31,34,37,40,43
Seq#:17; N Pos. 12,15,18,21,24,27,30,33,36
Seq#:18; N Pos. 10,13,16,19,22,25,28,31,34
Seq#:20; N Pos. 3,6,9,12,15,27,30,33,36,42
Seq#:21; N Pos. 7,13,16,19,22,31,37,40,43,46
Seq#:22; N Pos. 13,16,19,22,28,37,40,43,46,52
Seq#:23; N Pos. 19,25,28,31,34,40,43,46,55,61,64,67,70,76,79,82
Seq#:24; N Pos. 3,6,9,15,18,21,24,30,39,42,45,51,54,57,60,66
Seq#:26; N Pos. 7,13,16,19,22,25,31,34,43,49,52,55,58,64,67,70
Seq#:27; N Pos. 13,16,19,25,28,31,34,40,43,52,55,61,64,67,70,76
Seq#:28; N Pos. 19,25,28,31,34,40,43,46,55,61,64,67,70,76,79,82

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/804,733B

PAIR: 12182002

TIME: 12:14

Input Set : A:\MONS016US.TXT

Output Set: N:\CRF4\12182002\I804733B.raw

L:104 M:120 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
 L:105 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
 L:106 M:217 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
 L:107 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
 L:108 M:11 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6
 L:109 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
 L:110 M:7 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7
 L:111 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
 L:112 M:217 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8
 L:113 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
 L:114 M:11 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9
 L:115 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
 L:116 M:7 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:10
 L:117 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
 L:118 M:217 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:11
 L:119 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
 L:120 M:11 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12
 L:121 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
 L:122 M:7 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:13
 L:123 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0
 L:124 M:217 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14
 L:125 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0
 L:126 M:11 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:15
 L:127 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
 L:128 M:217 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:16
 L:129 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0
 L:130 M:11 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:17
 L:131 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
 L:132 M:217 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:18
 L:133 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
 L:134 M:11 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:19
 L:135 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
 L:136 M:217 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:20
 L:137 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
 L:138 M:11 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:21
 L:139 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
 L:140 M:217 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:22
 L:141 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
 L:142 M:11 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:23
 L:143 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
 L:144 M:217 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:24
 L:145 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
 L:146 M:11 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:25
 L:147 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0
 L:148 M:217 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:26
 L:149 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0
 L:150 M:11 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:27
 L:151 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
 L:152 M:217 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:28
 L:153 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/804,733B

DATE: 11/17/2002

TIME: 11:11:11

Input Set : A:\MONS016US.TXT

Output Set: N:\CRF4\12182002\I804733B.raw

L:664 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12
L:669 M:341 W: 146 "r" or "Naa" used, for SEQ ID#:12 after pos.:6
L:671 M:341 W: 146 "n" or "Naa" used, for SEQ ID#:12 after pos.:6
L:677 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12